

FIG. 1A

FIG. 1B

FIG. 1C

FIG. 1D

FIG. 1E

FIG. 1F

FIG. 1

1 GGATCCTGGT CGCGAGCGCG CCGCCCGGCC ACCTGCGGGC GCGCCCCGCC

GGGACCGCTC GAGGACGCGT CGCGAAGGCT CTAGGGGCTG TATCTTCAG

101 AGTCTACGCC CCTTTGTGCA GTGCACAAA TTTCCGTGCT AGCTTCATGC
"-35"

phaE
TATCACGCC CAGACGAGGA AGATTACCG TGAAACGATAC GGCCAACAAG
S/D V N D T A N K

201 ACCAGCGACT GGCTGGACAT CCAACCGCAAG TACTGGGAGA CCTGGTCGGA
T S D W L D I Q R K Y W E T W S E

GCTGGCGGC AAGACCTTGG GTCTGGAGAA GACCCCGGCC AATCCTTGGG
L G R E T L G L E K T P A N P W A

301 CGGGCGCCCT CGATCATTGG TGGCAGACGG TCTCGCCCGC CGCCCCCAAC
G A L D H W W Q T V S P A A P N

QACCTGGTTC CGGACTTCAT GGAGAAAGCTC GCGGAGGAGG GCAAGGCGTT
D I V R D I M E K I A F O G K A F

401 CTTGGCTC ACCGACTACT TCAugAUgg CCTCGGGGGGC AGTAGGUGA
F G E T D Y F T K G L G G S S G T

CGCAGGGCTG GGACACCCCTC TCGAAAGACCA TCGACGGACAT GCAAAGGCC

501 TTCGCCAGCG GCCGGATCGA AGGCGACGAG ACCTTCGGCC GCCTGATGGC
F A S G R I E G D E T F R R L M A
CTTCTGGGAG ATGCCGCTCG ACAACTGGCA GCGCACCATG TCCTCGCTGT
F W E M P L D N W Q R T M S S L S

601 CCCGGTGCC CGGCGACCTG CTGCGCAACA TGCCGCACGA CCAAGTCAGG
P V P G D L L R N M P H D Q V R
GACAGGGTCG ACCGCATCCT CTCGGCACCC GGGCTCGGCT ACACGCGCGA
D S V D R I L S A P G I G Y T R E

701 GGAGCAGGCC CGCTACCAGG ATCTGATCCG CCGCTCGCTG GAGTACCAAGT
E Q A R Y Q D L I R R S L E Y Q S
CGGCCCTGAA CGAATACAAC GGCTTCTTCG GCCAGCTCGG TGTCAAGTCC
A L N E Y N G F F G Q L G V K S

801 CTCGAGCGGA TGCGCGCCTT CCTGCAGGGA CAGGCCGAGA AGGGCGTCGC
L E R M R A F L Q G Q A E K G V A
CATCGAGTCG GCGCGCACCC TCTACCGACGC CTGGGTGGC TGCTGCGAAG
I F S A R T L Y D A W V G C C E E

901 AGTCATAGC CGAGGAGGTC AGCTCGGCG ACTACCGCGA CATCCACCGC
V Y A E E V S S A D Y A H E H G
CGCCTCGTCA ACGCCCAGAT GGCCCTCAAG CAGCGCATGT CGACCATGGT
R I V N A Q M A I K Q R M S I M V

FIG. 1B

1001 CGA(GAGGTC CTCGGCGCGA TGCGGCTGCG GACCCGCAGC GAGCTGCCA
 D E V L G A M P L P T R S E L R T
 CGC10CAGGA TCGGCTCCAG GAGTGGCGCG GCGAGGGCAA GCGCCAGCGC
 L Q D R L Q E S R G E G K R Q R

1101 CAAGAGATCG AGACGCTGAA GCGGCAGGTC GCGGCCTTGG CCGGCGGCCGC
 Q F I E T L K R Q V A A L A G G A
 CCAGCCCCGG CCCCAGGGCT CGGCCAGGCC CAGCACCCGG CCCGCGCCGG
 Q P A P Q A S A Q P S T R P A P A

1201 CGA(GGCCCG GGCGGCGAGC GCGGCGCCCA AGCGCAGCAC CACGACCCGC
 T A P A A S A A P K R S T T T R
 CGCAAGACCA CCAAGCCAC CACCGGCCAG TGATGTGGC CGCCCGTCCA
 R E T T K P T T G Q *

phaC →
 1301 TCGGCACCAAG GAGAGAGTGC CGTGTCGCCA TTCCCGATCG ACATCCGGCC
 S/D V S P F P I D I R P
 CGACAAGCTG ACCGGAGGAGA TGCTGGAGTA CAGCCGCAAG CTCGGCGAGG
 D K I F F I M L F Y S R K I G : G

1401 GIAIGAGAA CCTGGCTCAAG GGCGAUCAGA TCGACACAGG CGTCACCCCG
 M Q N L L K A D Q I D F G V T P
 AAGGAGGCTGG TCCACCGCGA GGACAAGCTG GTCCTCTACC GCTACCGGGG

1501 CCGGGCGCAG GTGGCGACCC AGACCGATCCC GCTGCTGATC GTCTACGCC
P A Q V A T Q T I P L L I V Y A L

TCGTCAATCG GCCCTACATG ACCGACATCC AGGAGGATCG CTGGACGATC
V N R P Y M T D I Q E D R S T I

1601 AAGGGCCTGC TCGCCACCGG TCAGGACGTC TATCTGATCG ACTGGGGCTA
K G L L A T G Q D V Y L I D W G Y

CCGGATCAG GCCGACCGGG CGCTGACCC CGATGACTAC ATCAACGGCT
P D Q A D R A I T L D D Y I N G Y

1701 ACATCGACCG CTGCGTCGAC TACCTGCGCG AGACCCACGG CGTCGACCAG
I D R C V D Y L R E T H G V D Q

GTCAACCTGC TCGGGATCTG CCAGGGCGGG GCCTTCAGCC TCTGCTACAC
V N L L G I C Q G G A F S L C Y T

1801 GGCCCTGCAC TCCGAGAAGG TCAAAAACCT CGTCACCATG GTCACGCCGG
A L H S E K V K N L V T M V T P V

TCGACTTCCA GACCCCCGGGC AACCTGCTCT CGGCCTGGGT CCAGAACGTC
D F Q T P G N I L S A W V Q N V

1901 GAGGTGAACT TGGGGCGGA CACCATGGC AACATCGG QUAATGCT
D V D E A V D I M G N I P G F T L

CAACTGGACC TTCCCTGTCGC TCAAGCCCTT CAGCCTGACC GGCCAGAAGT
N W F E L S I K P F S I T G Q K Y

FIG. 1D

2001 ACGTCAACAT GGTGACCTG CTCGACGACG AGGACAAGGT CAAGAACCTC
V N M V D L L D D E D K V R N F

CTGGGGATGG AGAAGTGGAT CTTCGACAGC CCGGACCAGG CGGGCGAGAC
L R M E K W I F D S P D Q A G E T

2101 CTTCCGCCAG TTCAATCAAGG ACTTCTACCA GCGCAACGGC TTCAATCAACG
F R Q F I K D F Y Q R N G F I N G

GCGGCGTCCT GATGGCGAT CAGGAGGTG ACCTGCGCAA CATCCGCTGC
G V I T G D Q E V D L R N I R C

2201 CGGGTCTGA ACATCTACCC GATGCAGGAC CACCTGGTGC CGCCGGATGC
P V L N I Y P M Q D H L V P P D A

CTCCAAGGCC CTGGCGGGAC TGACCTCCAG CGAGGACTAC ACGGAGCTCG
S K A L A G L T S S E D Y T E L A

2301 CCTTCCCCGG CGGGCACATC GGCAATCTACG TCAGCGGCAA GGCGCAGGAA
F P G G H I G I Y V S G K A Q E

GGAGTCACCC CGGGGATCGG CCGCTGGCTG AACGAACGGG GCTGAGCGGG
G V T P A T G R W L N E R G *

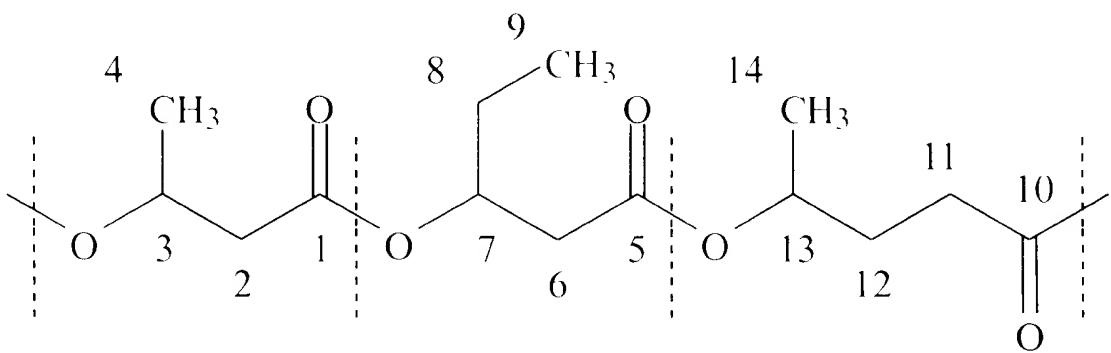
2401 GTCGACCCAC CGGGCTGGACG GGGCGGGCGG GGGCAATCGA AGGCGGGCGG
CGGGGGCGA TGGGGATGC GGGCGGGCGG CGGGGGGGG CGGACCGCG

2501 CCGCCGCACC CGCATCGCCC CGCGGGCTGG CGTACAATGA CGGTCTTCGC
GAGCGAGCCC CGCATCGTCA ACGGAGGGCTG CATGGGCGCC GACCACCAAAC

2601 TGCTGGCCGC GTACGACGGG CTGGCCGAGA CCTACGACGC CCACCGCGGC
CTCTTCGACA TGCGCGCCGT GCTCGAGGAC ATCTTCGCC GCCTGCCGGC

2701 CTGCGGCACC CTCTCGACC TCGGCTGCGG CGCCGGGGAG CCGTGCGCGC
GCGCCTTCCT CGACCGCGGC TGGCGGGTGA CCGGGGTGGA CTTCTGCCCG

2801 GCCATGCTCG CCCTCGCGGC GCGCTACGTC CCCGAGATGG AGCGGATCC



3HB

3HV

4HV

FIG. 2

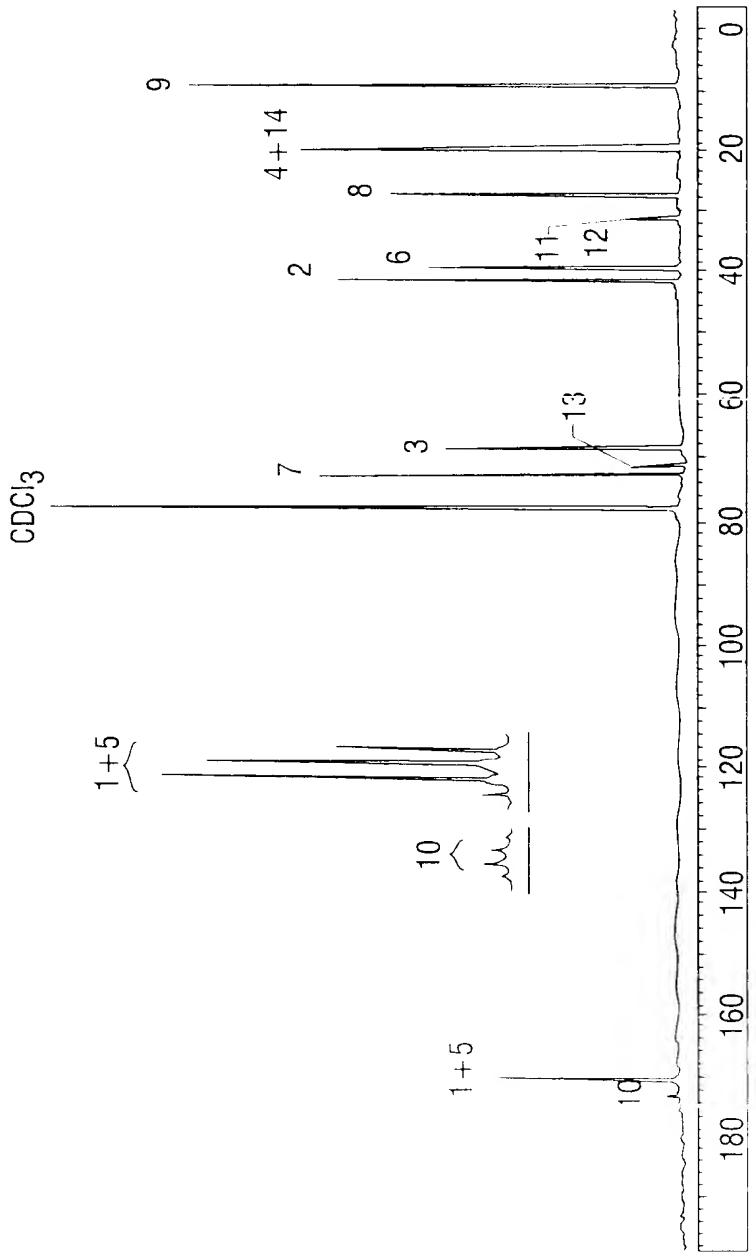


FIG. 3

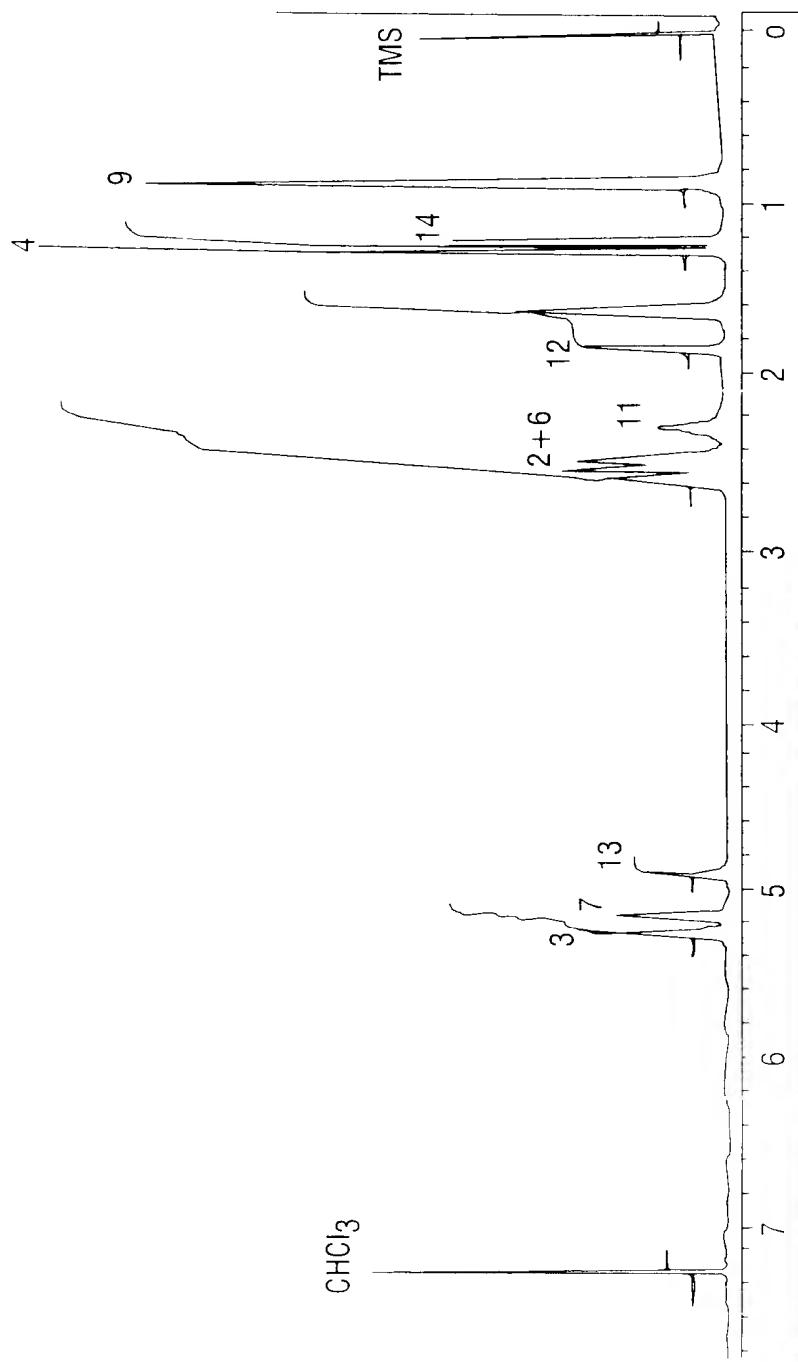


FIG. 4

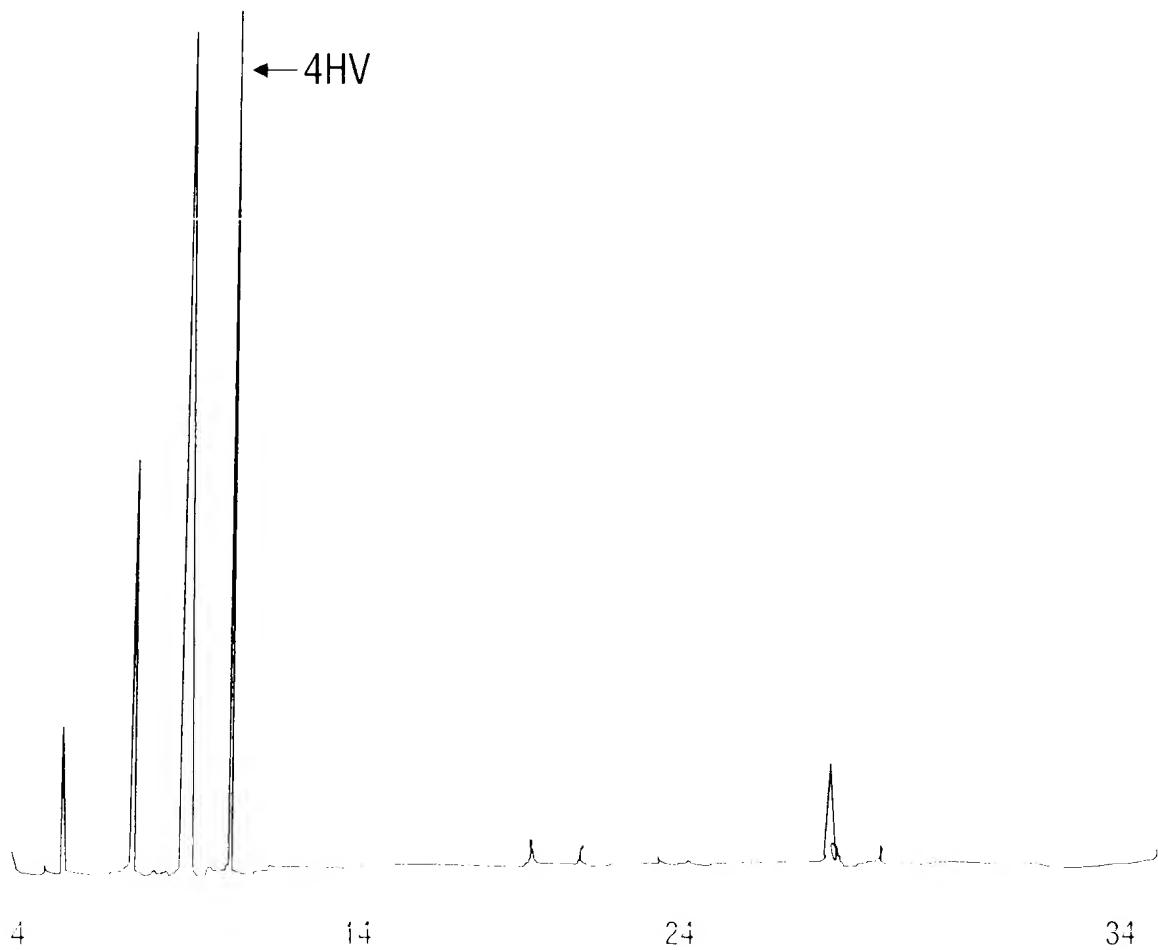


FIG. 5

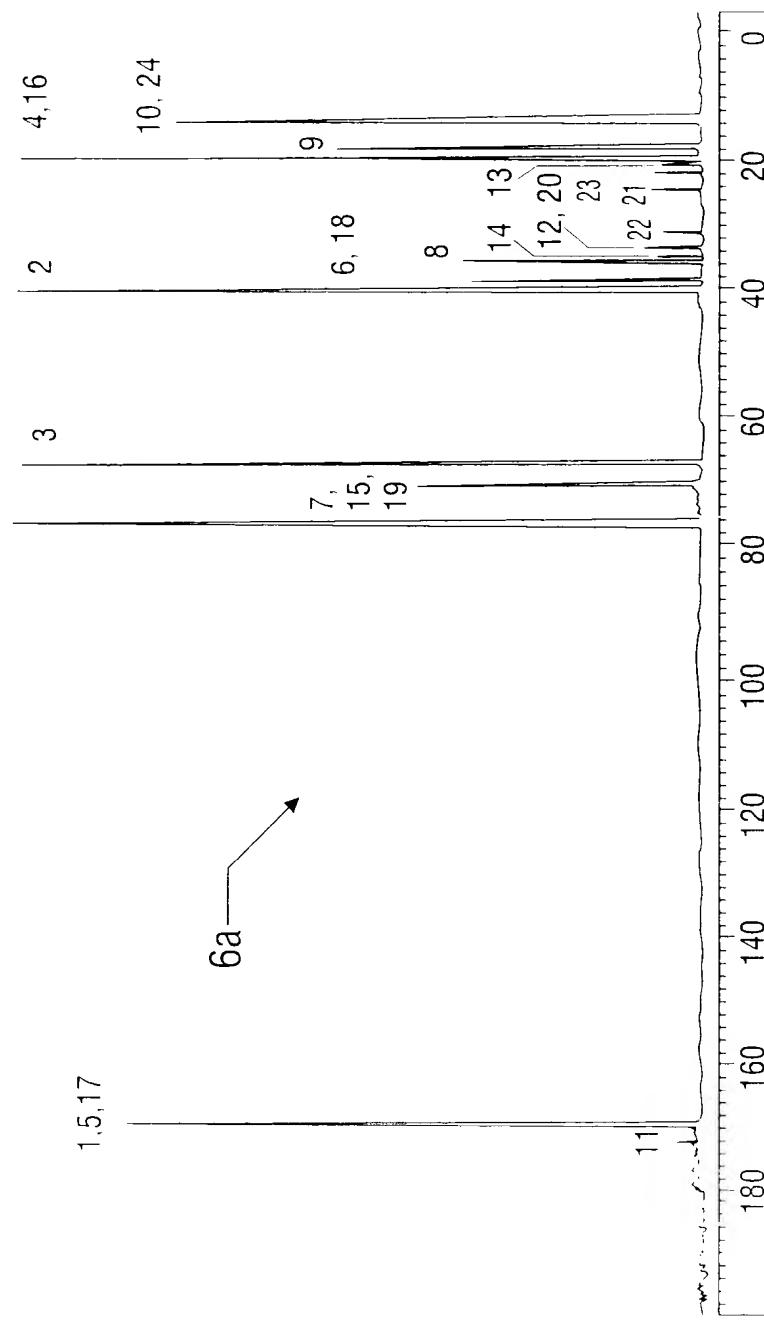
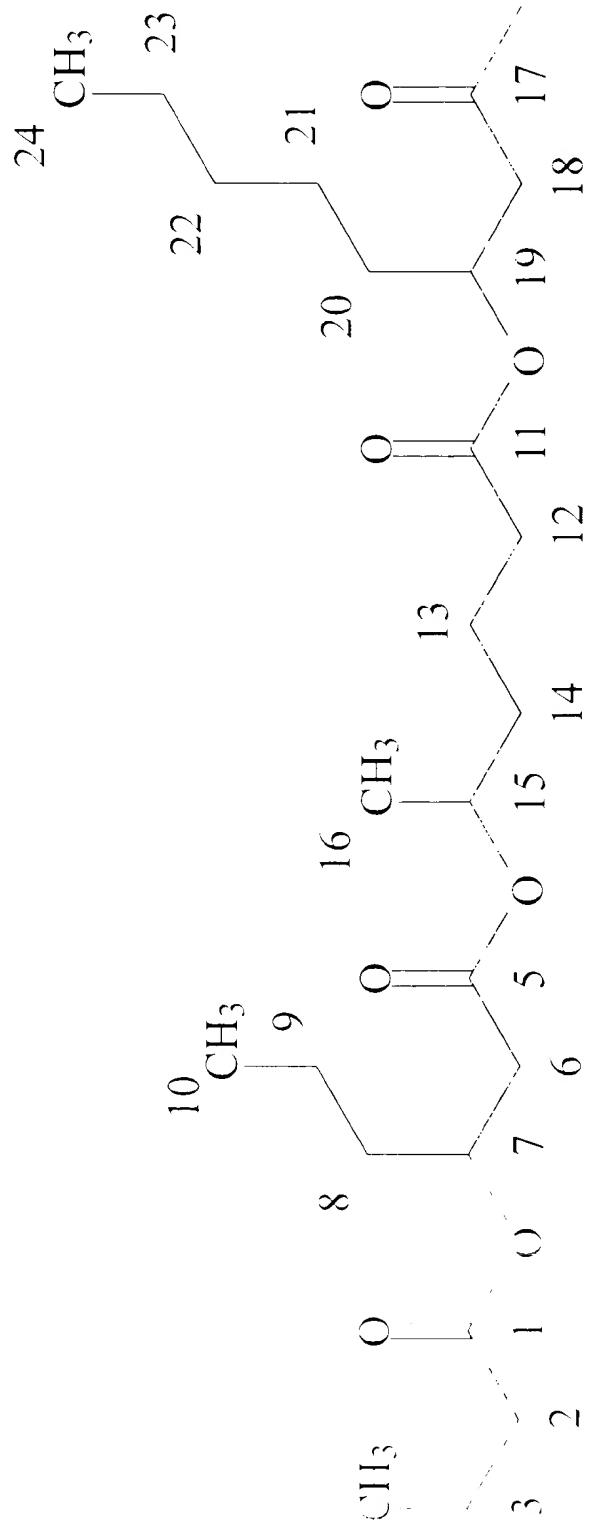


FIG. 6



3HO

5HHX

3HHX

3HB

FIG. 6A

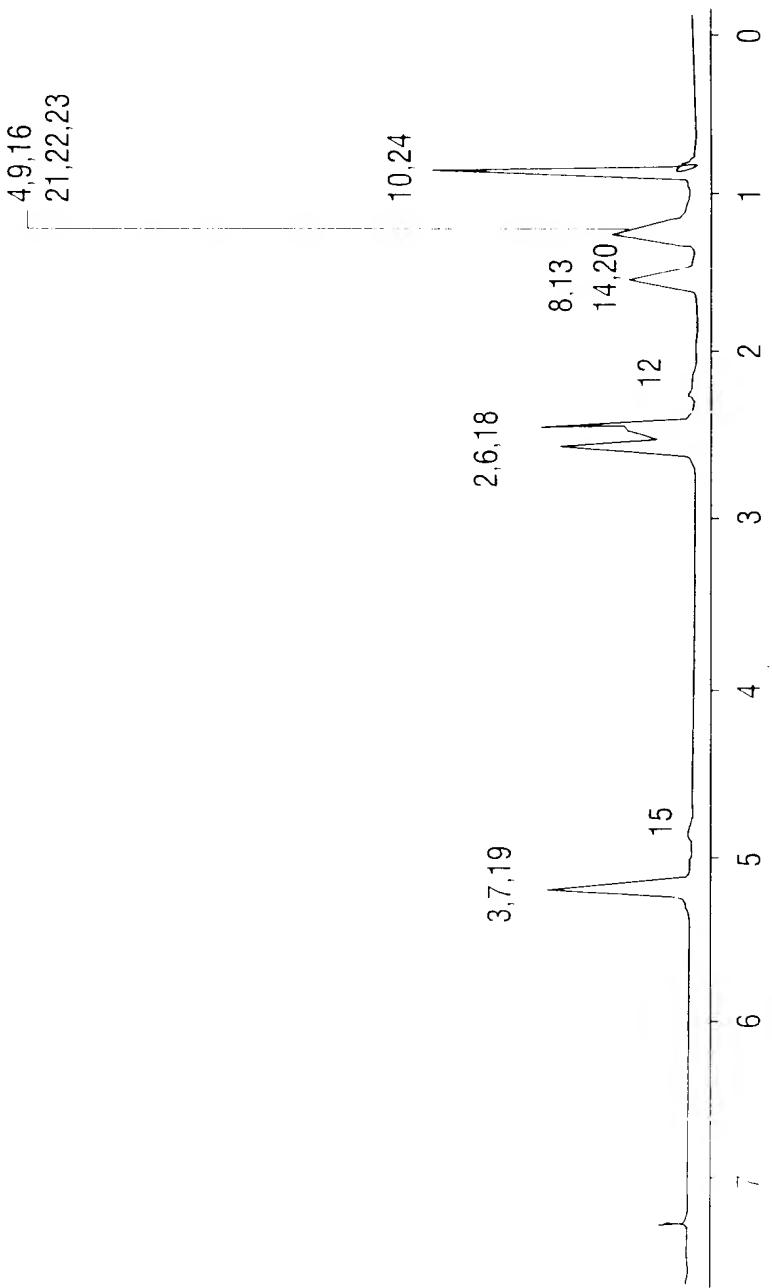


FIG. 7

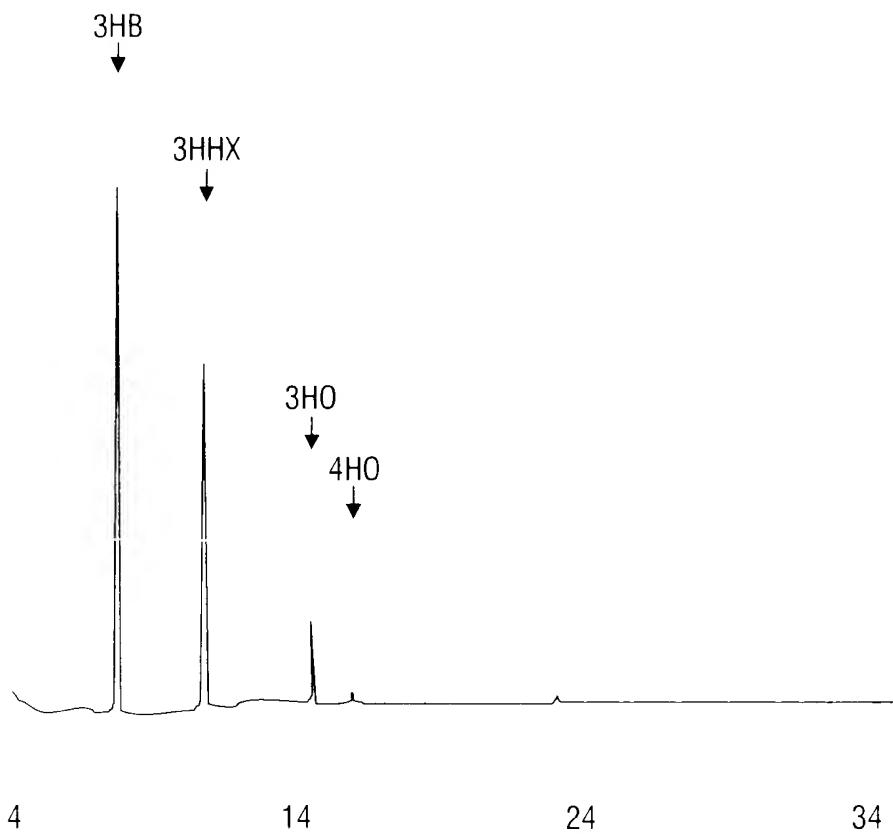


FIG. 8

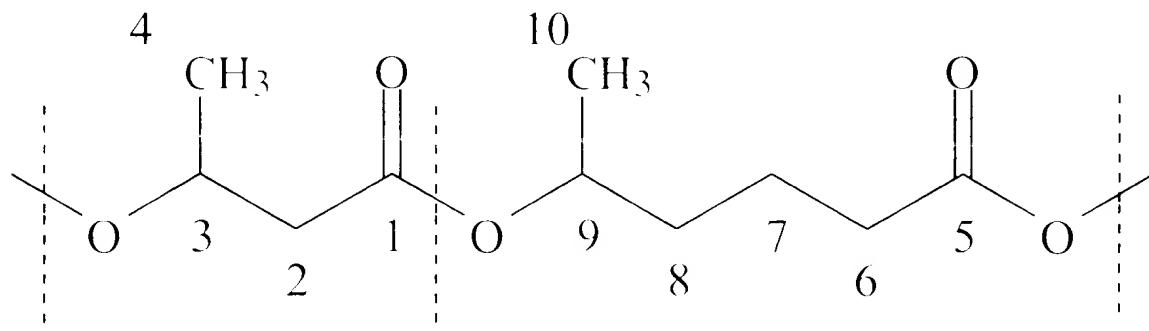


FIG. 9

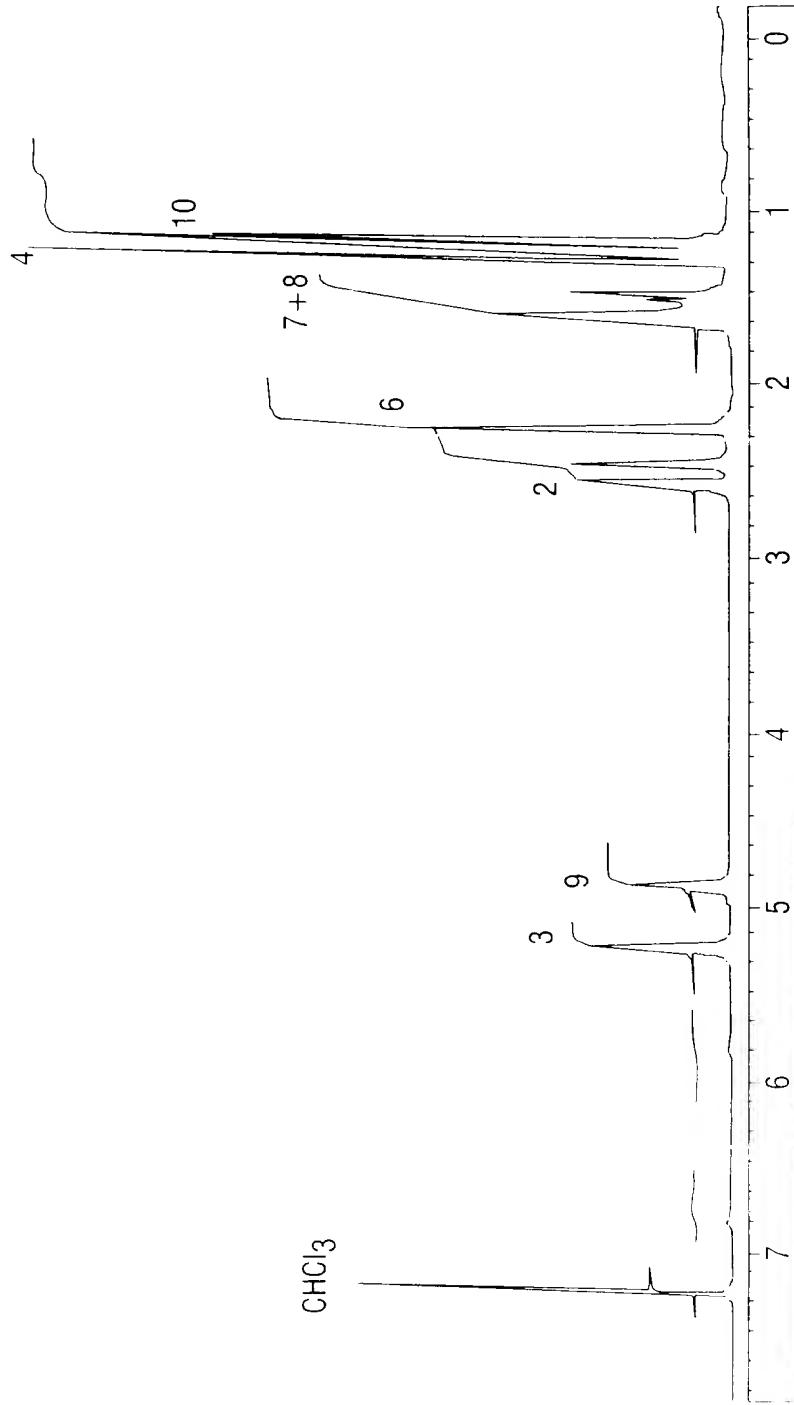


FIG. 10

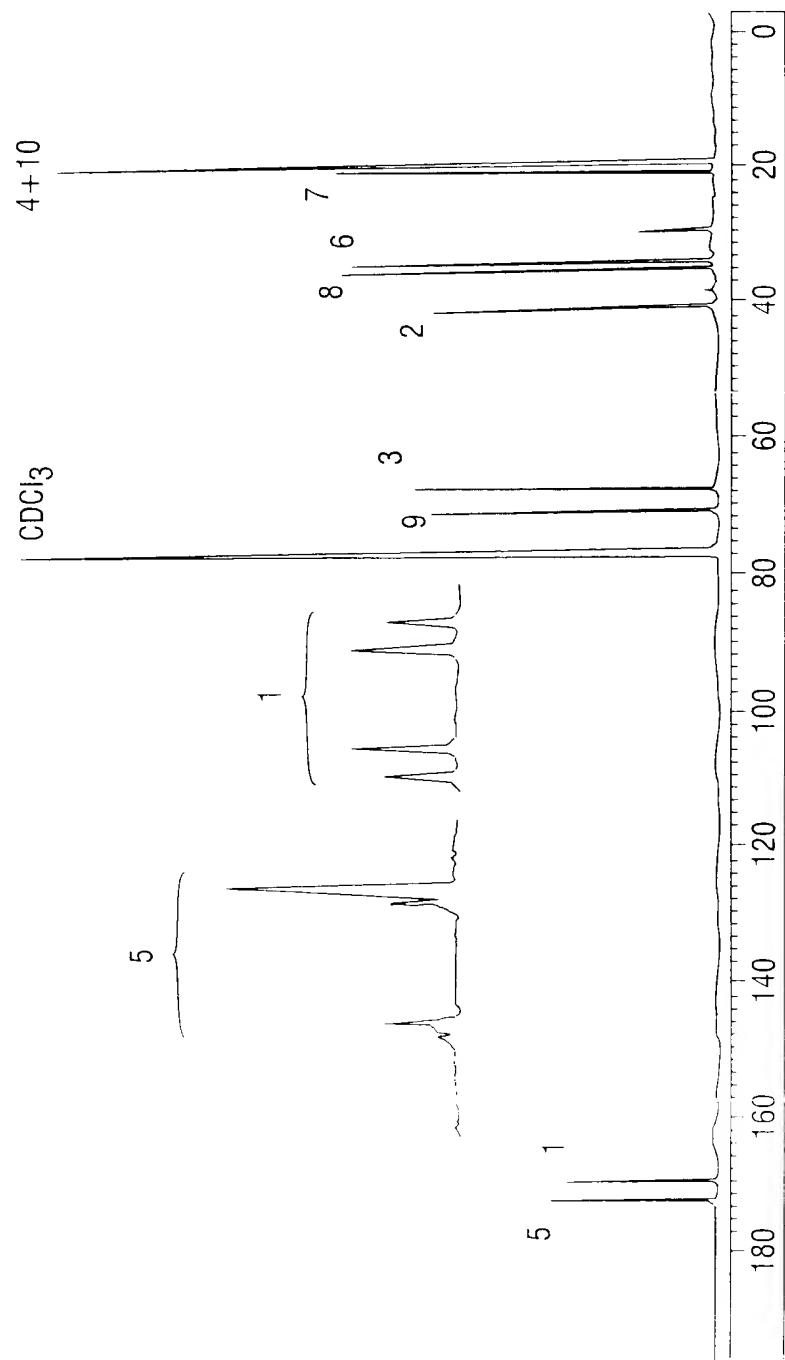


FIG. 11

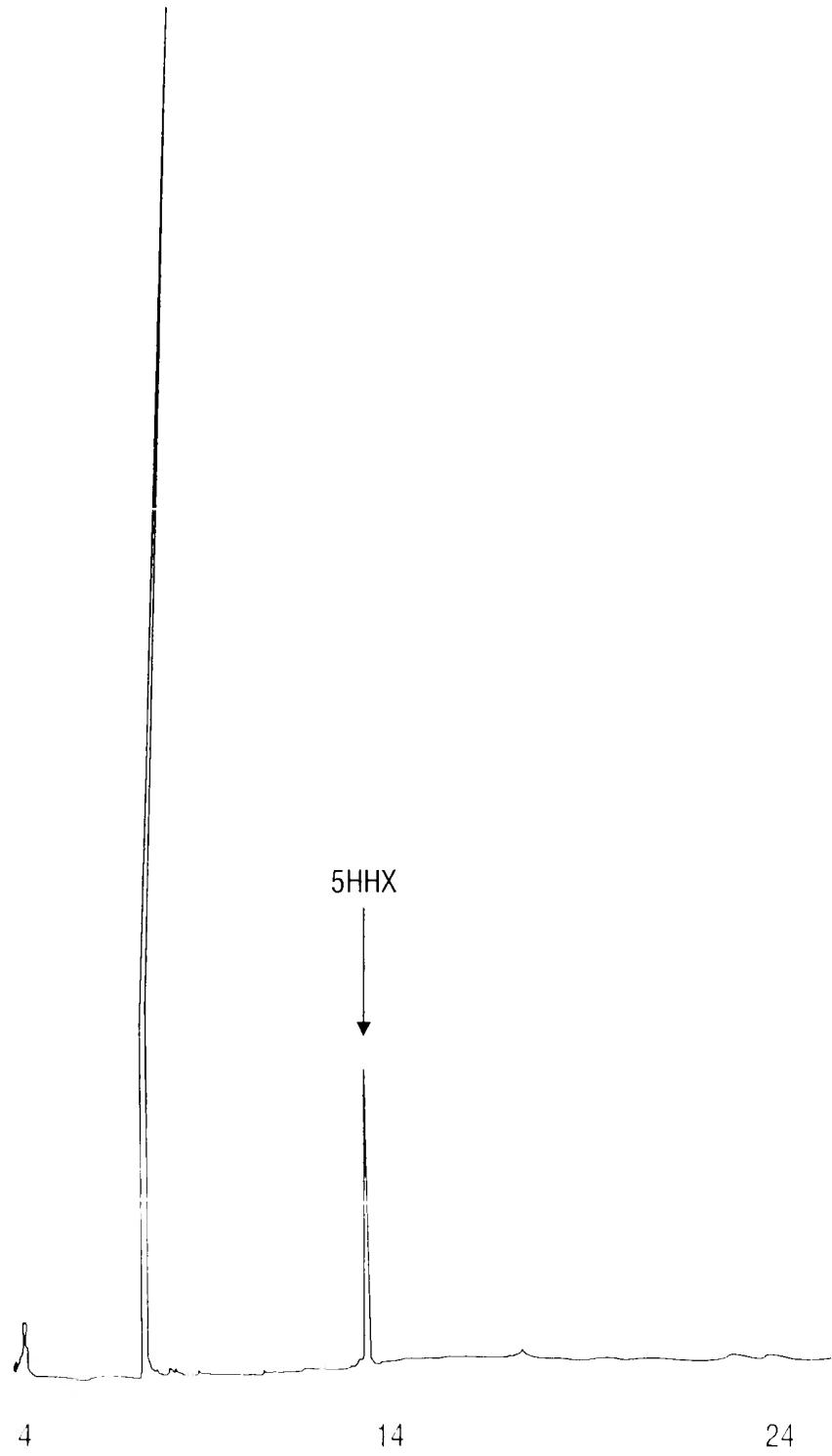
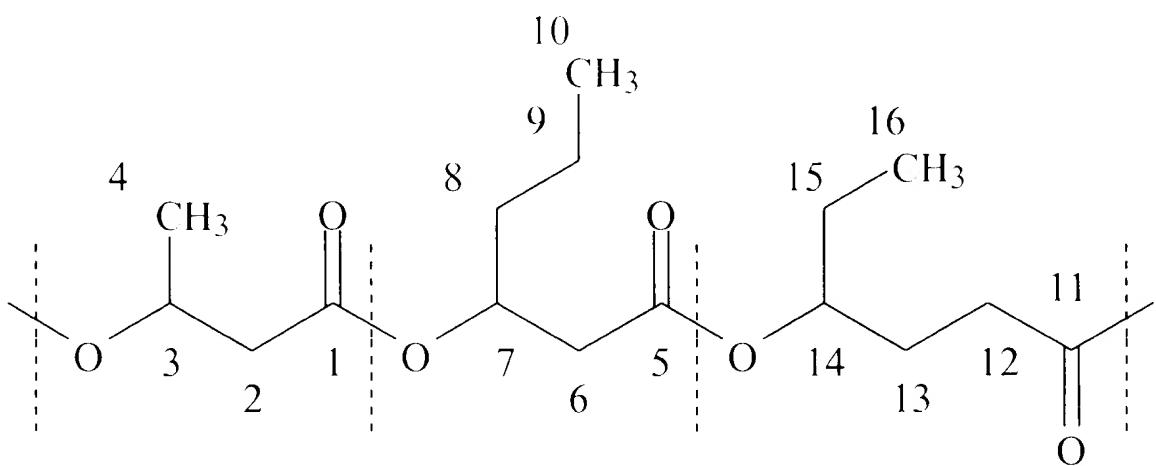


FIG. 12



3HB

3HHx

4HHx

FIG. 13

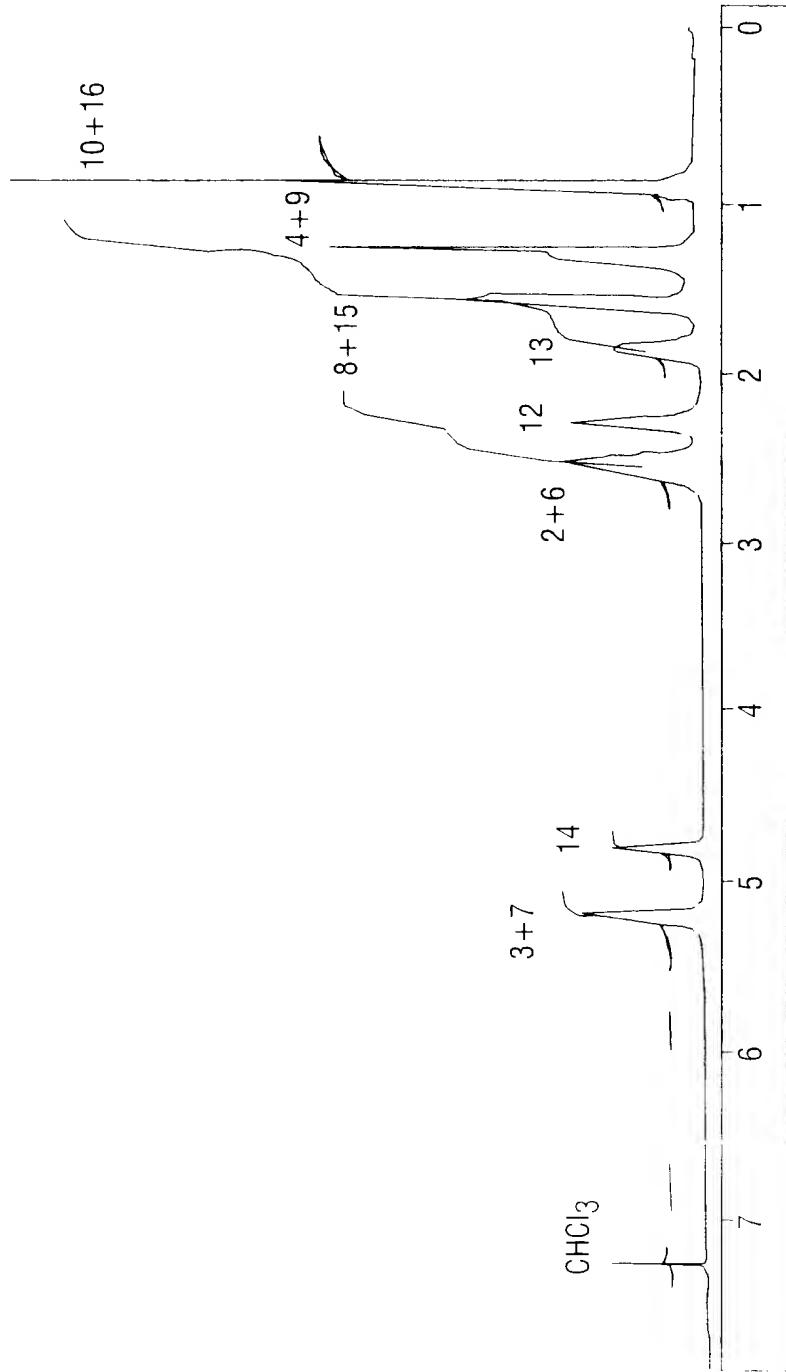


FIG. 14

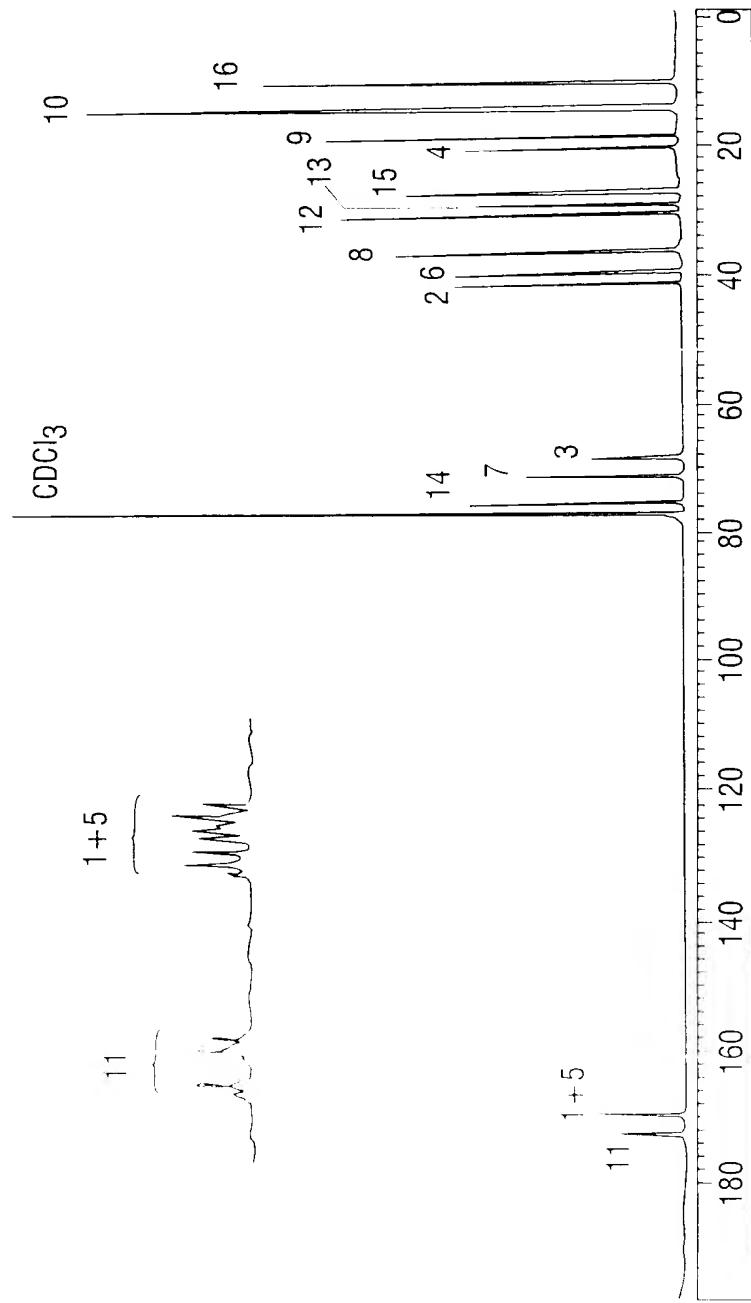


FIG. 15

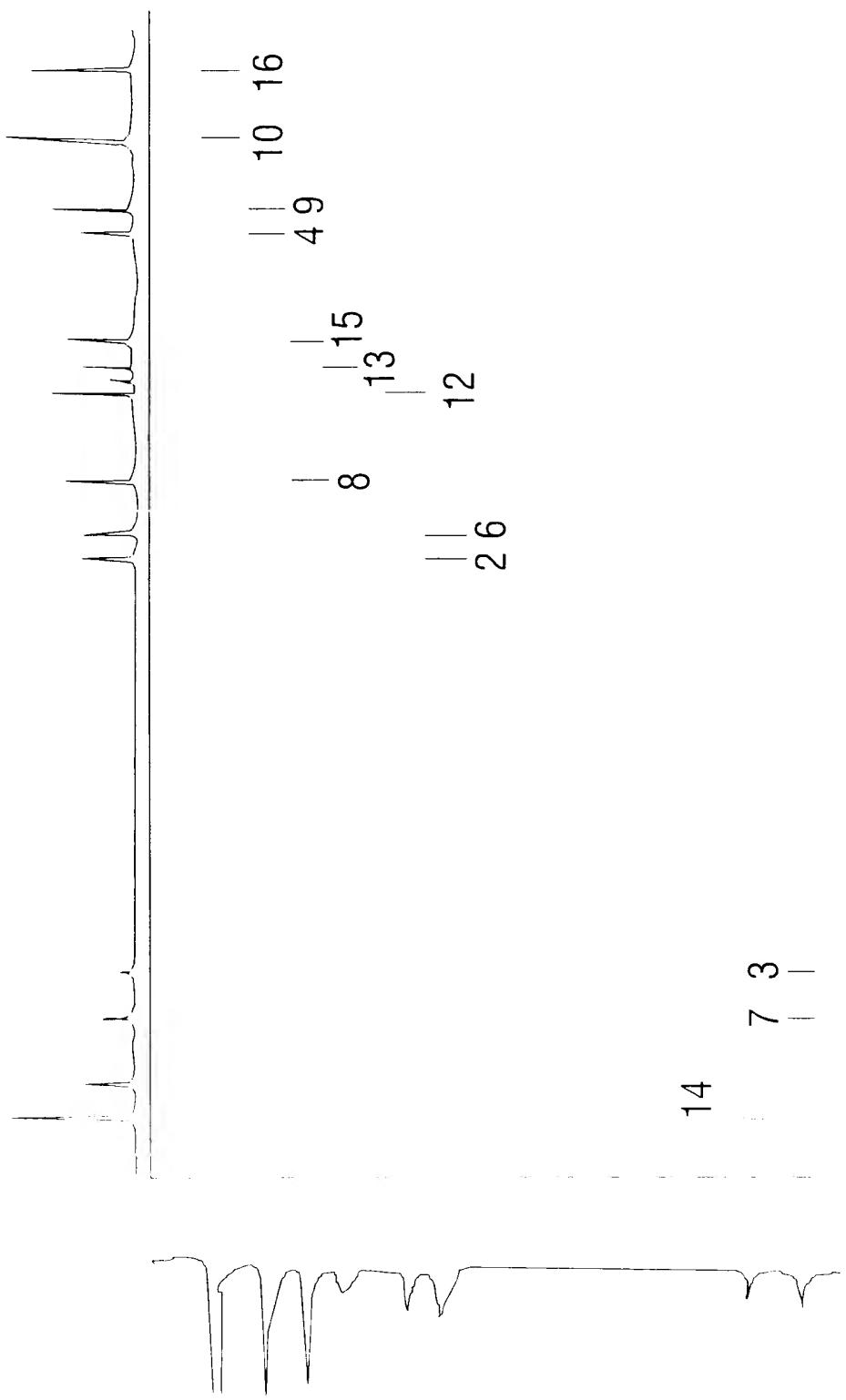


FIG. 16

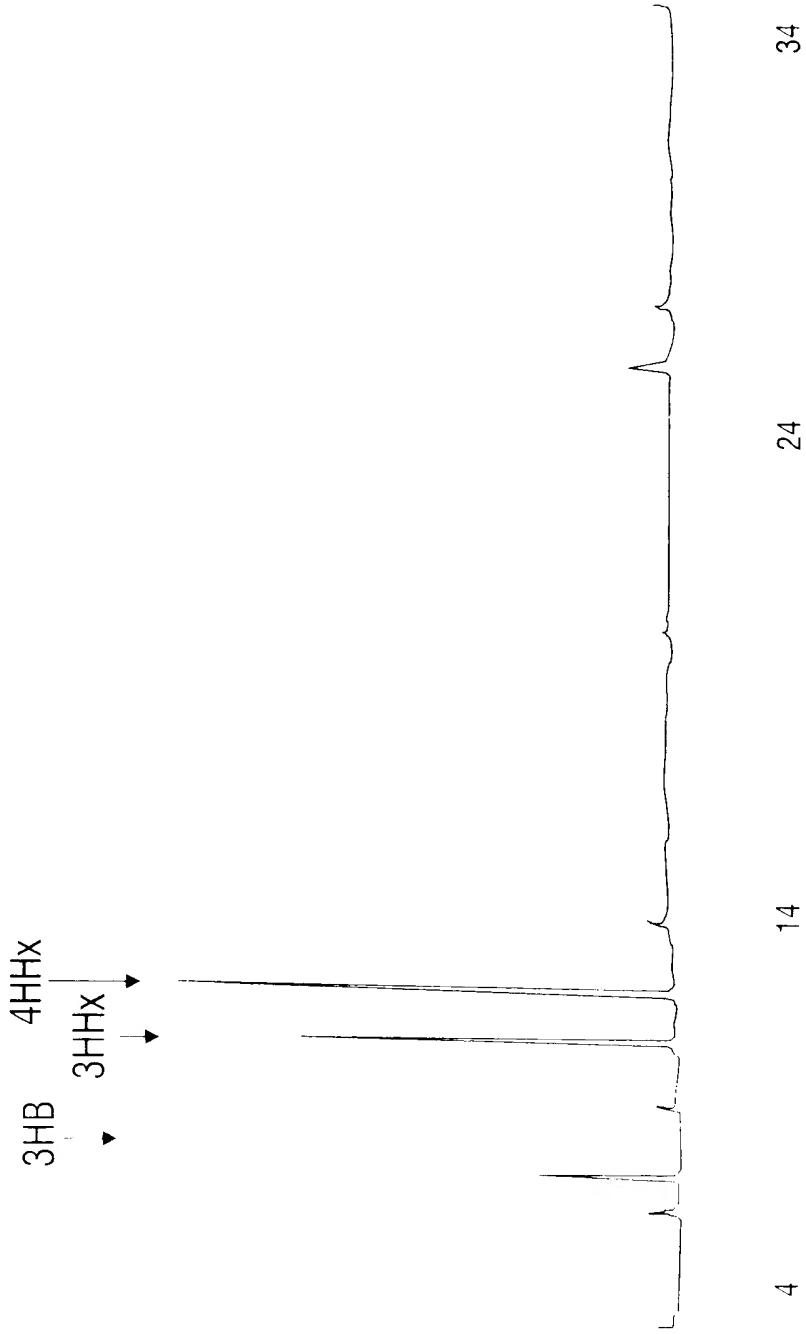


FIG. 17